SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT:
 - (A) NAME: ENFER TECHNOLOGY LIMITED ET AL
 - (B) STREET: EARLSFORT TERRACE
 - (C) CITY: DUBLIN
 - (E) COUNTRY: IRELAND
 - (F) POSTAL CODE (ZIP): 2
 - (G) TELEPHONE: 353-1-6605033
 - (H) TELEFAX: 353-1-6606920
- (ii) TITLE OF INVENTION: IMMUNOLOGICAL ASSAY FOR SPONGIFORM ENCE
- (iii) NUMBER OF SEQUENCES: 4
 - (iv) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)
 - (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: CHRISTINA GATES
 - (B) REFERENCE DOCKET NUMBER: PL678pct
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 353-1-6605033
 - (B) TELEFAX: 353-1-6606920
- (2) INFORMATION FOR SEQ ID NO: 1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 45
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: unknown
 - (ii) MOLECULE TYPE: Amino acid
 - (iii) HYPOTHETICAL:
 - (iv) ANTI-SENSE:
 - (vi) ORIGINAL SOURCE:
 - (A) ORGANISM:
 - (G) CELL TYPE:
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

Met Val Lys Ser His IIe Gly Ser Trp IIe Leu Val Leu
1 5 10

 Phe Val Val Ala Met Trp Ser Asp Val Gly Leu Cys Lys 20 Lys Arg Pro Lys Pro Gly Gly Gly Trp Asn Thr Gly Gly 30 Ser Arg Tyr Pro Gly Xaa

- (2) INFORMATION FOR SEQ ID NO: 2:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 43 base pairs
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: unknown
 - (ii) MOLECULE TYPE: Amino acid
 - (iii) HYPOTHETICAL:
 - (iv) ANTI-SENSE:
 - (vi) ORIGINAL SOURCE:
 - (A) ORGANISM:
 - (G) CELL TYPE:
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

W W W W GTy Ser Pro Gly Gly Asn Arg Tyr Pro Pro Gln Gly Gly Gly Trp Gly Gln Pro His Gly Gly Gly Trp Gly Gln . 20 Pro His Gly Gly Gly Trp Gly Gln Pro His Gly Gly Gly 35 Gin Gly Gln Pro 4.0.

- (3) INFORMATION FOR SEQ ID NO: 3:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 43 base pairs
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: unknown
 - (ii) MOLECULE TYPE: Amino acid
 - (iii) HYPOTHETICAL:
 - (iv) ANTI-SENSE:
 - (vi) ORIGINAL SOURCE:
 - (A) ORGANISM:
 - (G) CELL TYPE:
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

Gly Gly Gly Trp Gly Gln Gly Gly Ser His Ser Gln Trp Asn Lys Pro Ser Lys Pro Pro Lys Thr Asn Met 20 Lys His Val Ala Gly Ala Ala Gly Ala Val Gly Gly Leu Gly Gly Tyr 40

- (3) INFORMATION FOR SEQ ID NO: 4:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 46 base pairs
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: unknown
 - (ii) MOLECULE TYPE: amino acid
 - (iii) HYPOTHETICAL:
 - (iv) ANTI-SENSE:
 - (vi) ORIGINAL SOURCE:
 - (A) ORGANISM:
 - (G) CELL TYPE:
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

Met Leu Gly Ser Ala Met Ser Ser Pro Leu IIe His Phe Gy Asn Asp Tyr Glu Asp Arg Tyr Thr Arg Glu Asn Met 15 20 Tyr Arg Tyr Pro Asn Gln Val Tyr Tyr Arg Pro Val Asp 35

Tyr Ser Asn Gln Asn Asn